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Exhibit B

☐ **1:** NP_195636. Reports ...[gi:15234999] The record has been replaced by 145356202

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LOCUS NP_195636 97 aa linear PLN 09-JUN-2006
 DEFINITION DNA binding / transcription factor [Arabidopsis thaliana].
 ACCESSION NP_195636
 VERSION NP_195636.1 GI:15234999
 DBSOURCE REFSEQ: accession NM_120086.1
 KEYWORDS .
 SOURCE Arabidopsis thaliana (thale cress)
 ORGANISM Arabidopsis thaliana
 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
 Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
 rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
 COMMENT PROVISIONAL REFSEQ: This record has not yet been subject to final
 NCBI review. The reference sequence was derived from AT4G39250.1p.
 [WARNING] On Apr 18, 2007 this sequence was replaced by
 gi:145356202.
 Method: conceptual translation.
 FEATURES
 source Location/Qualifiers
 1..97
 /organism="Arabidopsis thaliana"
 /db_xref="taxon:3702"
 /chromosome="4"
 /ecotype="Columbia"
 Protein 1..97
 /product="DNA binding / transcription factor"
 /calculated_mol_wt=10697
 Region 14..60
 /region_name="SANT"
 /note="SWI3, ADA2, N-CoR and TFIIB' DNA-binding domains.
 Tandem copies of the domain bind telomeric DNA tandem
 repeats as part of the capping complex. Binding is sequence
 dependent for repeats which contain the G/C rich motif
 [C2-3 A (CA)1-6]; cd00167"
 Site /db_xref="CDD:29107"
 order(14,47..48,50..51,53..55,57..59)
 /site_type="DNA binding"
 /db_xref="CDD:29107"
 CDS 1..97
 /locus_tag="AT4G39250"
 /coded_by="NM_120086.1:1..294"
 /GO_component="chloroplast; nucleus"
 /GO_function="DNA binding; transcription factor activity
 [PMID 1118137]"
 /GO_process="regulation of transcription, DNA-dependent"
 /note="myb family transcription factor, contains PFAM
 profile: PF00249 myb-like DNA binding domain"
 /db_xref="GeneID:83081"
 /db_xref="TAIR:AT4G39250"
 ORIGIN
 1 massmsssq sgswtakgnk afeqalatyd qdtpnrqnv akvvggktte evkrhyellv

61 qdinsiengh vpfpnrtsg gctngrlsqe ekryvls

//

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Aug 28, 2012 16:53:42